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Responses of *Phanerochaete chrysosporium* to Toxic Pollutants: Physiological Flux, Oxidative Stress, and Detoxification

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Supporting Information

ABSTRACT: The white-rot fungus *Phanerochaete chrysosporium* has been widely used for the treatment of waste streams containing heavy metals and toxic organic pollutants. The development of fungal-based treatment technologies requires detailed knowledge of the relationship between bulk water quality and the physiological responses of fungi. A noninvasive microtest technique was used to quantify real-time changes in proton, oxygen, and cadmium ion fluxes following the exposure of *P. chrysosporium* to environmental toxic (2,4-dichlorophenol and cadmium). Significant changes in H⁺ and O₂ flux occurred after exposure to 10 mg/L 2,4-dichlorophenol and 0.1 mM cadmium. Cd²⁺ flux decreased with time. Reactive oxygen species formation and antioxidant levels increased after cadmium treatment. Superoxide dismutase activity correlated well with malondialdehyde levels (r² =



0.964) at low cadmium concentrations. However, this correlation diminished and malondialdehyde levels significantly increased at the highest cadmium concentration tested. Real-time microscale signatures of H⁺, O₂, and Cd²⁺ fluxes coupled with oxidative stress analysis can improve our understanding of the physiological responses of *P. chrysosporium* to toxic pollutants and provide useful information for the development of fungal-based technologies to improve the treatment of wastes cocontaminated with heavy metals and organic pollutants.

INTRODUCTION

As one of the most effective methods for environmental remediation, bioremediation has increasingly attracted the attention of environmentalists.¹ Bioremediation using microorganisms offers an attractive treatment option because this technology is cost-effective, environmentally compatible, and has high removal efficiency for various pollutants.^{1–3} A large number of microorganisms have been the focus of research for their potential pollutant disposal ability. White-rot fungi are the most efficient lignin degraders, and the representative species, *Phanerochaete chrysosporium*, has been extensively used for its ability to degrade a wide range of organic substrates.^{4–6} Generally, the organic substrates degraded by *P. chrysosporium* may coexist with heavy metals in wastewater, which can affect microbial reproduction and cause morphological and physiological changes.^{7,8}

The biodegradation or biosorption ability of microorganisms may deteriorate due to the morphological and physiological changes caused by heavy metals and toxic organic pollutants in the environment.^{9,10} For example, cadmium causes cellular toxicity via several mechanisms, including interference with DNA repair and protein metabolism, membrane lipid peroxidation, physiological Zn(II) substitution, and reactive oxygen species (ROS) formation.^{11,12} Oxygen free radicals, such as hydrogen peroxide (H_2O_2) , superoxide (O_2^-) , and hydroxyl (OH^-) radicals, are among the most reactive compounds induced by heavy metal stress.¹³ Excess free radicals could lead to multiple toxic effects, such as lipid peroxidation, protein cleavage, and DNA damage.^{14,15}

Removal of toxic pollutants from substrates is a challenge because they affect fungal colonization and bioactivity.¹⁶ It is necessary to understand the interactions between *P. chrysosporium* and the pollutants in the medium. Previous studies have shown that *P. chrysosporium* can tolerate heavy metals and toxic organic pollutants to some degree and remove them from wastewater.^{17–19} It is likely that multiple mechanisms, such as efflux pumps, modifying enzymes, target mutations, and resistance systems, are involved in microbial resistance to environmental stress (e.g., UV radiation or chemical toxic exposure).^{20–22} Therefore, *P. chrysosporium* can adapt to complex polluted environments.

Although P. chrysosporium studies on organic matter degradation and metal accumulation from liquid media can

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provide useful information on the relationship between fungal activities and pollutants,^{3,23} they do not accurately reflect the treatment process. Our previous study also demonstrated a decrease in the pH of the treatment system (from 6.5 to 3.6) and cadmium concentration-dependent alterations in protein production and enzymatic activities of P. chrysosporium;³ however, the underlying mechanisms remain unknown. To date, the effects of pollutant stress on the metabolism, resistance, and adaptive responses of P. chrysosporium have not been reported in the literature. Particularly, the effects of toxic pollutants on the physiological flux, oxidative stress, and detoxification of P. chrvsosporium have not been investigated. The relative toxicities and fates of pollutants occurring in the growth environment of P. chrysosporium remain unknown. From an environmental point of view, it is important to understand the responses of P. chrysosporium to toxic pollutants. Such information would enable the development of fungal-based technologies to improve the removal of metalpolluted organic wastes.

Therefore, the main purpose of this study was to examine the responses of *P. chrysosporium* to toxic pollutants using physiological measurements. We tested the toxic organic pollutant 2,4-dichlorophenol (2,4-DCP) and the heavy metal cadmium, which are typically found in sites cocontaminated with organic and heavy metal pollutants. To assess physiological responses, we monitored real-time changes in H⁺, O₂, and Cd²⁺ fluxes in *P. chrysosporium* following exposure to pollutants using noninvasive microtest technology. Further, antioxidative responses of *P. chrysosporium* to cadmium were also evaluated. These studies would help clarify the mechanisms involved in heavy-metal toxicity and/or elucidate the cellular basis of mass tolerance to these compounds.

MATERIALS AND METHODS

Strain and Chemicals. The *P. chrysosporium* strain BKMF-1767 (ATCC 24725) was obtained from the China Center for Type Culture Collection (Wuhan, China). Stock cultures were maintained on malt extract agar slants at 4 °C. Spores were gently scraped from the agar surface and blended in sterile distilled water to obtain a spore suspension. The spore concentration was adjusted to 2.0×10^6 spores/mL. Aqueous suspensions of fungal spores were inoculated into Kirk's liquid culture medium²⁴ in 500-mL Erlenmeyer flasks. All the chemicals used in this study were of analytical reagent grade.

Physiological Flux. After 3 days of growth in liquid medium, P. chrysosporium pellets were harvested and exposed to toxics. Net fluxes of H⁺, O₂, and Cd²⁺ were measured using the noninvasive microtest technique (the NMT system BIO-IM; YoungerUSA, LLC, Amherst, MA) and the Optical Oxygen Flux Measurement System (YGOO-01A; YoungerUSA) at Xuyue (Beijing) Science & Technology Co. Ltd., China (SI Figure S1). A schematic of P. chrysosporium H⁺, O₂, and Cd²⁺ flux measurements using microelectrodes is shown in SI Figure S2. The principle of NMT and its applications in ion flux detection have been described previously.²⁵⁻²⁷ Briefly, for H⁺ and Cd²⁺ flux measurements, prepulled and silanized glass micropipets (inner diameter, $3 \pm 1 \mu m$; XYPG120–2; Xuyue) were first filled with a backfilling solution (H⁺: 15 mM NaCl + 40 mM KH₂PO₄ at pH 7.0; Cd²⁺: 10 mM CdCl₂ + 0.1 mM KCl) to a distance of ~ 1 cm from the tip. The micropipets were then front-filled with approximately 30 μ m-columns of selective liquid ion-exchange cocktails (Hydrogen ionophore Icocktail B [Product No. 95293]; Cadmium Ionophore I,

[Product No. 20909]; Sigma-Aldrich, St Louis, MO). Ionselective electrodes were calibrated prior to flux measurements with different concentrations of target ion buffer (H⁺: pH 4.5 and 6.5; Cd²⁺: 0.05 and 0.5 mM). Only electrodes with Nernstian slopes greater than a certain value per decade (H⁺: 58 mV; Cd²⁺: 25 mV) were used in this study. The concentration gradients of the target ions were measured by moving the ion-selective microelectrode between 2 points close to the fungal pellet surface (ca. $3 \pm 1 \ \mu$ m) in a preset excursion with a distance of 30 μ m. During the measurement of Cd²⁺ flux, *P. chrysosporium* cells were treated with a Ca²⁺ channel inhibitor (GdCl₃, 0.1 mM), and changes in Cd²⁺ flux before and after the treatment were recorded.

For O₂ flux measurement, fiber-optic oxygen microsensors (optrodes) and a frequency-domain lifetime fluorometer were constructed using previously published techniques.²⁸ Optrodes (tip diameter, 5–7 μ m) were calibrated in sterile nitrogen-purged and O₂-saturated growth media (21%). The measured phase angle was transduced to an analog signal via a digital signal processor (YGOO-MC; YoungerUSA).^{29,30} The recording rate for the H⁺ and Cd²⁺ fluxes was 5.96 s/reading. For O₂, the entire cycle was completed in 8.91 s. Ionic/molecular fluxes were calculated using Fick's law of diffusion

$$J = -D\frac{dc}{dx}$$

where J represents the ion flux in the x direction, and dc/dx is the ion concentration gradient. In our experiments, dx was 30 μ m, which is the distance of the microelectrode movement between a close point and far point, and D is the ion diffusion coefficient (varying with the kind of ion) in a particular medium. Data and image acquisition, preliminary processing, control of the electrode position, and stepper motor-controlled fine focus of the microscope stage were performed with the imFlux software. All experiments were repeated 5 times to ensure the validity of physiological trends.

Antioxidant Enzymes. Superoxide dismutase (SOD) activity was measured according to the method reported by Choudhary et al.³¹ The cells were harvested by centrifugation and homogenized in 0.05 M phosphate buffer (pH 7.8). The homogenate was centrifuged at 15,000 rpm at 4 °C, and the supernatant was used for enzyme assay. SOD activity was assayed by monitoring the inhibition of photochemical reduction of nitroblue tetrazolium chloride (NBT) in a reaction mixture containing 1 M Na₂CO₃, 200 mM methionine, 2.25 mM NBT, 3 mM EDTA, 60 mM riboflavin, and 0.1 M phosphate buffer (pH 7.8). Absorbance was read at 560 nm.

Catalase (CAT) activity was measured in a 3-mL reaction mixture containing 50 μ L of enzymatic extract, 20 mM H₂O₂, and 50 mM potassium phosphate buffer (pH 7.0), according to the method reported by Cavalcanti et al.³² The disappearance of H₂O₂ was evaluated by measuring the decrease in absorbance at 240 nm (molar extinction coefficient ε = 36.6 mM⁻¹ cm⁻¹).

Lipid Peroxidation and Superoxide (O₂⁻) Production. Lipid peroxidation was evaluated by measuring the concentration of malondialdehyde (MDA), which is a cytotoxic product of lipid peroxidation and an indicator of free radical production and consequent tissue damage.³¹ MDA content was estimated using the procedure described by Aravind and Prasad,³³ with minor modifications. The harvested *P. chrysosporium* cells were homogenized in 10% trichloroacetic acid and centrifuged at 10,000 rpm for 15 min. The supernatant was boiled with thiobarbituric acid for 20 min. The heated supernatant was centrifuged at 5000 rpm for 5 min, and the absorbance was measured at 532 and 600 nm using the UV–visible spectrometer, Specord 200 PC.

 O_2^- production was analyzed as described by Lei et al.³⁴ The harvested cells were homogenized in 2 mL of 50 mM potassium phosphate buffer (pH 7.8) and centrifuged at 10,000 rpm for 10 min at 4 °C. One milliliter of the supernatant was mixed with 0.9 mL of 50 mM potassium phosphate buffer (pH 7.8) and 0.1 mL of 10 mM hydroxylamine hydrochloride. The reaction mixture was incubated at 25 °C for 20 min before adding 1 mL of 17 mM p-aminobenzenesulfonic acid and 1 mL of 7 mM α -naphthylamine. After further incubation (25 °C, 20 min), the absorbance of the mixture was spectrophotometrically recorded at 530 nm.

Statistical Analysis. The mean changes in H⁺, O₂, and Cd²⁺ fluxes were calculated by averaging continuous flux measurements during pre-exposure and peak exposure. The results are presented as the mean of 5 replicates, and standard deviations were used to analyze experimental data. A one-way analysis of variance (ANOVA) was used to test for significant differences ($\alpha < 0.05$) between the flux values for each treatment. All reported error bars represent one standard error of the arithmetic mean.

RESULTS

Chemical Toxic Exposure and Stress Response. Representative real-time plots of *P. chrysosporium* H^+ and O_2 fluxes during exposure to 2,4-DCP and cadmium are presented in Figure 1A (for the average value around the fungal surface, see Figure 1B). After exposure to 2,4-DCP (10 mg/L), the H⁺ influx increased compared with the pre-exposure value (preexposure: -33 ± 4 pmol cm⁻² s⁻¹; peak stress response: $-37 \pm$ 3 pmol cm^{-2} s⁻¹; throughout the manuscript, negative and positive values represent influx and efflux, respectively). The oxygen flux of P. chrysosporium initially increased upon exposure to 10 mg/L 2,4-DCP (4 ± 0.5 min) and subsequently decreased; the stable O2 influx of P. chrysosporium after exposure to 2,4-DCP was higher than the pre-exposure value. The stable O₂ flux values during pre-exposure and exposure were -44 ± 0.5 and -45 ± 1 pmol cm⁻² s⁻¹, respectively. These results were different from those for Nitrosomonas europaea biofilm during chlorocarbonyl cyanide phenylhydrazone exposure.²² After the subsequent addition of 0.1 mM Cd(NO₃)₂, the H⁺ flux changed from influx to efflux (13 \pm 5 pmol cm⁻² s⁻¹). The O₂ efflux rapidly decreased, indicated by the high slope of the O₂ influx plot in Figure 1A, after the addition of 0.1 mM Cd(NO₃)₂. The monitored real-time H⁺ and O2 fluxes following sequential exposure P. chrysosporium to 2,4-DCP or Cd²⁺ only are presented in SI Figures S3-S5, and the results are described in the Supporting Information.

The Cd²⁺ flux of *P. chrysosporium* after exposure to 0.1 mM Cd(NO₃)₂ is shown in Figure 2. An obvious Cd²⁺ influx (-66 \pm 2 pmol cm⁻² s⁻¹) into *P. chrysosporium* pellets was observed in the initial 15 \pm 0.2 min of exposure. The following two reasons may account for this phenomenon: adsorption of Cd²⁺ by *P. chrysosporium* mycelia and entry into the cells following penetration of the cell wall and plasma membrane. Subsequent addition of 0.1 mM GdCl₃ (Ca²⁺ channel inhibitor)³⁵⁻³⁷ elicited a dramatic decrease in Cd²⁺ influx from -66 \pm 2 pmol cm⁻² s⁻¹ to -45 \pm 3 pmol cm⁻² s⁻¹, indicating that Ca²⁺ channels were involved in Cd²⁺ influx.

Figure 3 demonstrates the Cd^{2+} flux of *P. chrysosporium* in response to different durations of 0.1 and 0.5 mM $Cd(NO_3)_2$



Figure 1. Proton and oxygen fluxes of *P. chrysosporium* during exposure to 10 mg/L 2,4-DCP and 0.1 mM $Cd(NO_3)_2$. (A) Representative real-time plots. (B) Average values at the fungal surface. "Pre" means pre-exposure.



Figure 2. Real-time Cd^{2+} flux of *P. chrysosporium* during exposure to 0.1 mM $Cd(NO_3)_2$ and 0.1 mM $GdCl_3$.

treatment. The Cd²⁺ flux of *P. chrysosporium* following treatment with 0.1 mM Cd(NO₃)₂ at 0, 1, 3, and 6 h were -65 ± 4 , -20 ± 1 , -6 ± 0.2 , and 0.2 ± 0.7 pmol cm⁻² s⁻¹, respectively. The Cd²⁺ influx decreased with the duration of exposure, which reflects a gradual saturation of the adsorption and uptake processes. A similar trend was observed when 0.5 mM Cd(NO₃)₂ was used. The Cd²⁺ flux exhibited a small efflux (0.2 ± 0.7 pmol cm⁻² s⁻¹) and a small influx (-0.1 ± 1.5 pmol cm⁻² s⁻¹) at the 6 h time point of treatment with 0.1 mM and



Figure 3. Cd^{2+} fluxes of *P. chrysosporium* in response to different durations of treatment with 0.1 and 0.5 mM $Cd(NO_3)_2$. Numbers on the abscissa represent the exposure time. "0 h" indicates the measurement of Cd^{2+} flux immediately after exposure.

0.5 mM Cd(NO₃)₂, suggesting that equilibrium was achieved and the Cd²⁺ flux was in a state of dynamic equilibrium (SI Figures S6 and S7). Similar results were found when studying BY-2 cells Cd²⁺ flux following exposure to 150 μ M Cd²⁺.³⁸ In the present study, 0.5 mM Cd(NO₃)₂ elicited a Cd²⁺ influx of -121 ± 4 pmol cm⁻² s⁻¹, which was higher than that following treatment with 0.1 mM Cd(NO₃)₂. However, when the *P. chrysosporium* pellets were exposed to 0.5 mM Cd(NO₃)₂ for 1 h, the Cd²⁺ influx dramatically decreased to -5 ± 1 pmol cm⁻² s⁻¹, which was lower than that following a 1-h treatment with 0.1 mM Cd(NO₃)₂. This may be attributed to different driving forces and Cd²⁺ toxicity in *P. chrysosporium*.

The percent changes in flux of *P. chrysosporium* pellets following exposure to 2,4-DCP and $Cd(NO_3)_2$ are shown in Figures S8 and 4. All physiological stress responses were



Figure 4. Average Cd^{2+} flux and change in Cd^{2+} flux of *P. chrysosporium* during exposure to 0.1 mM $Cd(NO_3)_2$ and 0.1 mM $GdCl_3$.

significantly different from those of null controls. After the addition of 2,4-DCP, O₂ and H⁺ fluxes increased by 3% and 6% respectively, relative to pre-exposure levels. Upon addition of 0.1 mM Cd(NO₃)₂, the H⁺ flux shifted from influx (-37 ± 3 pmol cm⁻² s⁻¹) to efflux (13 ± 5 pmol cm⁻² s⁻¹). The average oxygen flux decreased by 9% in the initial 10 \pm 0.5 min of exposure. A larger reduction ($30 \pm 1\%$) in Cd²⁺ influx was observed following treatment with 0.1 mM gadolinium (GdCl₃), which is known to block stretch-activated calcium

channels, indicating the involvement of extracellular cadmium influx via these channels.

Cd²⁺-Induced Oxidative Stress and Detoxification. The biological responses to cadmium exposure are presented in Figure 5. Enzymatic activities and ROS levels were



Figure 5. Biological responses of *P. chrysosporium* to various concentrations of $Cd(NO_3)_2$. A 3-day-old culture was treated with 0, 0.01, 0.05, 0.1, or 0.5 mM $Cd(NO_3)_2$ solution and further cultivated for 2 h. The cells were then collected and washed twice with distilled water for the above analysis.

significantly altered in the cadmium-treated group compared to control (untreated group). SOD activity and MDA content increased in a concentration-dependent manner after exposure to cadmium solution. Another antioxidant enzyme, CAT, exhibited a bell-shaped response with a maximum activity at 0.1 mM Cd(NO₃)₂. A similar trend was also observed for O_2^{-1} generation. MDA content increased with increasing cadmium concentration in the culture medium, indicating concentrationdependent free radical generation. The SOD activity and MDA levels were well correlated $(r^2 = 0.951 \text{ and } r^2 = 0.990,$ respectively) with lower cadmium concentrations (0.01, 0.05, and 0.1 mM). However, the correlation coefficient decreased when the higher cadmium concentration (0.5 mM) was taken into account ($r^2 = 0.462$ and $r^2 = 0.808$) (Figure 6). For low cadmium concentration treatment (0.01 Mm-0.1 mM), the CAT activity and O_2^- were correlated ($r^2 = 0.971$ and $r^2 =$ 0.837, respectively) with cadmium concentrations. CAT activity was also well correlated with O_2^{-} levels ($r^2 = 0.814$) in this case (SI Figures S10 and S11). However, correlations between P. chrysosporium CAT activity, O_2^- levels, and $Cd(NO_3)_2$ concentration were not observed when the higher cadmium concentration (0.5 mM) was taken into account, because the CAT activity and O_2^- levels of *P. chrysosporium* exposed to 0.5 mM Cd(NO3)2 were much lower than those of P. chrysosporium exposed to 0.1 mM $Cd(NO_3)_2$.

DISCUSSION

Microbial metabolism is complex, and cells have a wide array of physiological defense mechanisms for survival (e.g., efflux pumps, facultative electron transport, resistance systems).³⁹ Fungi are more resistant to chemical stress following acclimation, and species such as *P. chrysosporium* have the ability to survive in the presence of certain chemical toxic.^{16,40} The growth environment of *P. chrysosporium* is often complex and toxic. Thus, detailed characterization of the interactions



Figure 6. Correlation between *P. chrysosporium* superoxide dismutase (SOD) activity, malondialdehyde (MDA) level, and $Cd(NO_3)_2$ concentration in the medium.

between *P. chrysosporium* and toxic pollutants would provide useful information for the development of fungal-based technologies to improve the degradation of metal-polluted organic wastes.

The dynamics of homeostatic H⁺ transport/cytoplasmic pH regulation in microorganisms depends on many factors (e.g., local microenvironment, species type, mode of energy transport), and oxygen transport is complicated by facultative respiration.⁴¹ The increased H⁺ influx in *P. chrysosporium* was a response to 2,4-DCP-induced oxidative stress. The presence of this toxic pollutant in the medium may alter the bioactivity of the cell or perturb the structure of membrane phospholipids, thereby increasing H⁺ permeability across the cytoplasmic membrane.^{42,43} The increased O_2 influx following the addition of 10 mg/L 2,4-DCP is indicative of a possible improvement in respiration. Aromatic compounds can be used as carbon and energy sources for the removal of heavy metals from a medium containing microorganisms.² Our previous study also showed that a low concentration of 2,4-DCP (<20 mg/L) in the medium is beneficial for cadmium removal.³ Thus, the 2,4-DCP (10 mg/L) in this study may be used as a carbon and energy source to improve respiration and increase O₂ influx. The H⁺ flux changed from influx to efflux after the addition of 0.1 mM $Cd(NO_3)_2$ to the medium. This may be due to the removal of Cd^{2+} from the medium via the bonding of Cd^{2+} to the multiple functional groups on the P. chrysosporium mycelial cell wall surface (e.g., -OH, -COOH, and -NH₂). The bonding of Cd^{2+} to these groups would release H^+ , thereby increasing H^+ efflux. Cd^{2+} in the medium may enter the *P. chrysosporium* cell during the uptake process and boost the fungal production of organic acids, such as oxalic acid,⁴⁴ which may further increase H^+ efflux.

The rapid decrease in O_2 flux following the addition of 0.1 mM $Cd(NO_3)_2$ is indicative of a significant decrease in viability. Cadmium is the most toxic heavy metal for all white-rot fungi and can affect many vital processes. For example, it inhibits growth and reduces fresh biomass, alters morphology, and stimulates the activity of antioxidant enzymes.^{45,46} It is wellknown that cadmium has a high affinity for the sulfhydryl groups of proteins and thereby inhibits SH-bearing, redoxregulated enzymes in many cellular processes.^{47,48} Cadmium also binds to and activates calmodulin and plays an important role in Ca²⁺-dependent regulatory pathways. It may affect cell wall production by inhibiting extracellular Ca²⁺ influx, promoting intracellular Ca^{2+} efflux, and disrupting the cytoplasmic Ca^{2+} gradient.⁴⁹ Cadmium entry into *P*. chrysosporium during the uptake process affects both individual reactions and complex metabolic processes, including inhibition of growth and protein synthesis.⁷ Moreover, Cd²⁺ may cause various types of cellular damage (e.g., Ca2+ release, ROS production, and DNA damage) leading to apoptosis.^{8,50,51}

A larger reduction $(30 \pm 1\%)$ in Cd²⁺ influx was observed following treatment with 0.1 mM GdCl₃ (Figures 2 and 5), which may be due to shutdown of the uptake pathway by the added GdCl₃. In research studies on cellular responses to mechanical stimuli, the cell membrane is a major target for the external mechanical forces that act upon a cell. Ion channels play a crucial role in the ionic permeability of biological membranes and are present in all cells tested so far.⁵² Ion channels detect and transduce external mechanical forces into electrical and/or chemical intracellular signals.53 Gadolinium ion (Gd^{3+}) , which is a lanthanide, plays an important role in the structure and function of biomembranes. Furthermore, in the ion channels research field, Gd^{3+} is a well-known inhibitor of mechanosensitive ion channels.^{37,54} The presence of submillimolar concentrations of gadolinium (Gd^{3+}) in the medium is sufficient to provoke the near-complete inhibition of shockinduced efflux of metabolites such as lactose and ATP in Escherichia coli and yeast and ATP in Streptococcus faecalis.55 Further, in patch-clamp experiments, gadolinium inhibits the giant stretch-activated channels of E. coli, S. faecalis, and Bacillus subtilis. All these data suggest that stretch-activated channels are localized in the cytoplasmic membrane, where they control metabolite flux and play a major role in the response to environmental stress.⁵⁶ The presence of gadolinium (Gd^{3+}) can also directly inhibit Ca²⁺ influx via fusion, which disrupts the Ca²⁺ gradient and causes cellular damage.³⁷ Thus, the added GdCl₃ may decrease Cd²⁺ influx by shutting down the Cd²⁺ uptake pathway (Ca²⁺ channel) and disrupting the Ca²⁺ gradient in P. chrysosporium cells.

Figure 3 shows the Cd^{2+} flux of *P. chrysosporium* exposed to 0.1 and 0.5 mM $Cd(NO_3)_2$ for different durations. A high concentration of cadmium enhances the driving force for mass transfer to overcome the mass transfer limitation between the *P. chrysosporium* pellets and the fluid phase, thereby increasing the Cd^{2+} influx.⁵⁷ In addition, an elevated cadmium concentration increases the number of collisions between the pellets and Cd^{2+} , thereby enhancing the uptake process.^{58,59} The enhanced uptake reduces the time required to achieve saturation. Thus, following exposure to 0.5 mM $Cd(NO_3)_2$ for

1 h, the Cd²⁺ influx dramatically decreased from -121 ± 4 pmol cm⁻² s⁻¹ to -5 ± 1 pmol cm⁻² s⁻¹, which was lower than that following a 1-h treatment with 0.1 mM Cd(NO₃)₂. Moreover, a high Cd²⁺ concentration in the medium was likely to have caused extensive damage to the *P. chrysosporium* cells, leading to a decrease in cellular bioactivity.

In our study, enzymatic activities and ROS levels significantly responded to metal exposure. However, the concentrationresponse relationships were not always monotonic, and, in 2 cases (CAT and O_2^{-}), bell-shaped concentration-response curves were noted: a significant increase was observed in the lower concentration range (<0.1 mM) and a decline was observed at higher metal concentrations (>0.1 mM). Free radical generation in P. chrysosporium, evaluated by MDA production, increased under heavy metal stress (Figure 5), which is similar to the effect of heavy metals in higher plants.^{32,60} This suggests that the toxic effect of heavy metals may be exerted via free radical generation. Heavy metals induce the production of ROS, including superoxide radical (O_2^{-}) , hydrogen peroxide (H_2O_2) , and hydroxyl radical (OH^-) and produce a variety of damaging effects collectively termed as oxidative stress.⁶¹ Under normal circumstances, cells can reduce oxygen to water via their electron transport chains and protect themselves from ROS-induced damage by using enzymes such as SOD (to convert superoxide to hydrogen peroxide) and CAT (to convert hydrogen peroxide to water and oxygen).⁶² Under unfavorable conditions such as heavy metal exposure, oxidative stress occurs and the accumulated ROS can rapidly attack biomolecules, including nucleic acids, proteins, lipids, and amino acids, leading to irreparable metabolic dysfunction and cell death.^{31,61}

MDA measurement is used as an indicator of lipid peroxidation, which is linked to the production of $O_2^{-.60}$ The elevated MDA levels suggest that metal ions stimulate the free radical-generating capacity of the microorganism. The generation of cadmium-induced ROS will stimulate the production of antioxidant enzymes to protect the cellular components from damage. Thus, the increase in SOD activity observed in our study is a response to the elevated MDA levels. SOD activity was well correlated with MDA levels $(r^2 = 0.964)$ at low cadmium concentrations (0.01 mM to 0.1 mM). However, when the ROS levels exceed the ability of the antioxidant system to cope with them, cellular damage occurs, indicated by the reduced correlation between SOD activity and MDA levels $(r^2 = 0.873)$ (SI Figure S9). Exposure to 0.5 mM Cd(NO₃)₂ did not elicit an obvious increase in SOD activity, and the CAT activity was lower than that of 0.1 mM Cd(NO₃)₂. These observations are consistent with the findings of Lichtenthaler⁶ who described the stress responses of plants as bell-shaped curves and considered the decreasing part as an exhaustion phase in which defense systems are overloaded, leading to chronic damage and cell death. Similarly, the nonlinear and bell-shaped curves observed in this study at high concentrations may reflect cytotoxicity due to ROS overproduction induced by the interactions between excess metals and cellular components. The excess superoxide radical disrupts signaling pathways that activate genes encoding antioxidant enzymes, such as SOD. Similar results were obtained by Dazy et al.⁶⁴ for Fontinalis antipyretica Hedw. Although genome/proteome damage was not quantified in the present study, the observed relationships between cellular damage (MDA and O_2^{-} level) and antioxidant enzyme levels suggest that the tolerance of P.

chrysosporium to heavy metals partially depends on its ability to prevent oxidative stress.

Physiological changes are often the earliest events in cellular stress response (e.g., changes in respiration/growth rate/ homeostasis), and many of these mechanisms are used by cells to protect the genome/proteome from damage.²² Results from this study will serve as useful references for the physiological responses of fungi to environmental toxic exposure. Further studies combining genomic/proteomic analysis and patch-clamp measurements are required to understand whether changes in physiological transport are linked to global changes in cell viability (genetic damage and mutation) and transmembrane signaling.

ASSOCIATED CONTENT

Supporting Information

More details of the noninvasive microtest system; real-time and average changes in H⁺, O₂, and Cd²⁺ flux at different treatments; correlations among SOD activity, MDA content, CAT activity, O₂⁻ levels, and Cd(NO₃)₂ concentration in the medium. This material is available free of charge via the Internet at http://pubs.acs.org.

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Notes

The authors declare no competing financial interest.

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